## SEQUENCE LISTING

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<110> Kwon, Byoung S.
<120> MURINE 4-1BB GENE
<130> 740.009US1
<140> US 08/012,269
<141> 1993-02-01
<150> US 07/922,996
<151> 1992-07-30
<150> US 07/267,572
<151> 1988-11-07
<160> 13
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 2350
<212> DNA
<213> Mus musculus
<220>
<221> misc_feature
<222> (1)...(2350)
\langle 223 \rangle n = A,T,C or G
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ctacaccagg aaaaggacac attcgacaac aggaaaggag cctgtcacag aaaaccacag
                                                                       180
tgtcctgtgc atgtgacatt tcgccatggg aaacaactgt tacaacgtgg tggtcattgt
                                                                       240
gctgctgcta gtgggctgtg agaaggtggg agccgtgcag aactcctgtg ataactgtca
                                                                       300
gcctggtact ttctgcagaa aatacaatcc agtctgcaag agctgccctc caagtacctt
                                                                       360
ctccagcata ggtggacagc cgaactgtaa catctgcaga gtgtgtgcag gctatttcag
                                                                       420
gttcaagaag ttttgctcct ctacccacaa cgcggagtgt gagtgcattg aaggattcca
                                                                       480
ttgcttgggg ccacagtgca ccagatgtga aaaggactgc aggcctggcc aggagctaac
                                                                       540
gaagcagggt tgcaaaacct gtagcttggg aacatttaat gaccagaacg gtactggcgt
                                                                       600
ctgtcgaccc tggacgaact gctctctaga cggaaggtct gtgcttaaga ccgggaccac
                                                                       660
ggagaaggac gtggtgtgtg gacccctgt ggtgagcttc tctcccagta ccaccatttc
                                                                       720
tgtgactcca gagggaggac caggagggca ctccttgcag gtccttacct tgttcctggc
                                                                       780
gctgacatcg gctttgctgc tggccctgat cttcattact ctcctgttct ctgtgctcaa
                                                                       840
atggatcagg aaaaaattcc cccacatatt caagcaacca tttaagaaga ccactggagc
                                                                       900
ageteaagag gaagatgett gtagetgeeg atgteeacag gaagaagaag gaggaggagg
                                                                       960
aggetatgag etgtgatgta etateetagg agatgtgtgg geegaaaceg agaageacta
                                                                      1020
ggaccccacc atcctgtgga acagcacaag caaccccacc accctgttct tacacatcat
                                                                      1080
cctagatgat gtgtgggcgc gcacctcatc caagtctctt ctaacgctaa catatttgtc
                                                                      1140
tttacctttt ttaaatcttt ttttaaattt aaattttatg tgtgtgagtg ttttgcctgc
                                                                      1200
ctgtatgcac acgtgtgtgt gtgtgtgtgt gtgacactcc tgatgcctga ggaggtcaga
                                                                      1260
agagaaaggg ttggttccat aagaactgga gttatggatg gctgtgagcc ggnnngatag
                                                                      1320
gtcgggacgg agacctgtct tcttatttta acgtgactgt ataataaaaa aaaaatgata
                                                                      1380
tttcgggaat tgtagagatt ctcctgacac ccttctagtt aatgatctaa gaggaattgt
                                                                      1440
tgatacgtag tatactgtat atgtgtatgt atatgtatat gtatatata gactctttta
                                                                      1500
ctgtcaaagt caacctagag tgtctggtta ccaggtcaat tttattggac attttacgtc
                                                                      1560
acacacacac acacacac acacacacgt ttatactacg tactgttatc ggtattctac
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gtcatataat gggatagggt aaaaggaaac caaagagtga gtgatattat tgtggaggtg

1620

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1680
acagactacc ccttctgggt acgtagggac agacctcctt cggactgtct aaaactcccc
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ttagaagtct cgtcaagttc ccggacgaag aggacagagg agacacagtc cgaaaagtta
                                                                      1800
tttttccggc aaatcctttc cctgtttcgt gacactccac cccttgtgga cacttgagtg
                                                                      1860
tcatccttgc gccggaaggt caggtggtac ccgtctgtag gggcggggag acagagccgc
                                                                      1920
gggggagcta cgagaatcga ctcacagggc gccccgggct tcgcaaatga aacttttta
                                                                      1980
atctcacaag tttcgtccgg gctcggcgga cctatggcgt cgatccttat taccttatcc
                                                                      2040
tggcgccaag ataaaacaac caaaagcctt gactccggta ctaattctcc ctgccggccc
                                                                      2100
ccgtaagcat aacgcggcga tctccacttt aagaacctgg ccgcgttctg cctggtctcg
                                                                      2160
ctttcgtaaa cggttcttac aaaagtaatt agttcttgct ttcagcctcc aagcttctgc
                                                                      2220
tagtctatgg cagcatcaag gctggtattt gctacggctg accgctacgc cgccgcaata
                                                                      2280
agggtactgg gcggcccgtc gaaggccctt tggtttcaga aacccaaggc cccctcata
                                                                      2340
ccaacgtttc gactttgatt cttgccggta cgtggtggtg ggtgccttag ctctttctcg
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<210> 2 <211> 256 <212> PRT

<213> Mus musculus

<400> 2

Met Gly A	sn Asn Cys 5	Tyr Asn	Val Val	Val Ile 10	Val Leu	Leu Leu 15	Val
Gly Cys G	lu Lys Val 20	Gly Ala	Val Gln 25	Asn Ser	Cys Asp	Asn Cys	Gln
_	hr Phe Cys 5	Arg Lys	Tyr Asn 40	Pro Val	Cys Lys 45	Ser Cys	Pro
Pro Ser T 50	hr Phe Ser	Ser Ile 55	Gly Gly	Gln Pro	Asn Cys	Asn Ile	Cys
Arg Val C	ys Ala Gly	Tyr Phe	Arg Phe	Lys Lys 75	Phe Cys	Ser Ser	Thr 80
His Asn A	la Glu Cys 85	Glu Cys	Ile Glu	Gly Phe 90	His Cys	Leu Gly 95	Pro
Gln Cys T	hr Arg Cys 100	Glu Lys	Asp Cys 105	<del>-</del>	Gly Gln	Glu Leu 110	Thr
_	ly Cys Lys 15	Thr Cys	Ser Leu 120	Gly Thr	Phe Asn 125	Asp Gln	Asn
Gly Thr G	ly Val Cys	Arg Pro 135	_	Asn Cys	Ser Leu 140	Asp Gly	Arg
	eu Lys Thr	_	Thr Glu	_	Val Val	<del>-</del> /	Pro 160
Pro Val V	al Ser Phe 165		Ser Thr	Thr Ile 170	Ser Val	Thr Pro 175	Glu
Gly Gly P	ro Gly Gly 180	His Ser	Leu Gln 185	Val Leu	Thr Leu	Phe Leu 190	Ala
	er Ala Leu 95	Leu Leu	Ala Leu 200	Ile Phe	Ile Thr 205	Leu Leu	Phe
Ser Val L 210	eu Lys Trp	Ile Arg 215	<del>-</del>	Phe Pro	His Ile 220	Phe Lys	Gln
Pro Phe L 225	ys Lys Thr	Thr Gly 230	Ala Ala	Gln Glu 235	Glu Asp	Ala Cys	Ser 240
Cys Arg C	ys Pro Gln 245		Glu Gly	Gly Gly 250	Gly Gly	Tyr Glu 255	Leu

<210> 3

<211> 24

<212> PRT

<213> Mus musculus

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<400> 3
Cys Arg Val Cys Ala Gly Tyr Phe Arg Phe Lys Lys Phe Cys Ser Ser
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Thr His Asn Ala Glu Cys Glu Cys
            20
<210> 4
<211> 22
<212> PRT
<213> Drosophila
<400> 4
Cys Pro Val Cys Phe Asp Tyr Val Ile Leu Gln Cys Ser Ser Gly His
 1
                 5
                                                          15
Leu Val Cys Val Ser Cys
            20
<210> 5
<211> 26
<212> PRT
<213> Dictyostelium
<400> 5
Cys Pro Ile Cys Phe Glu Phe Ile Tyr Lys Lys Gln Ile Tyr Gln Cys
                                                          15
Lys Ser Gly His His Ala Cys Lys Glu Cys
            20
                                 25
<210> 6
<211> 6
<212> PRT
<213> Mus musculus
<220>
<221> SITE
<222> (1)...(6)
<223> Xaa = Any Amino Acid
<400> 6
Val Gln Asn Ser Xaa Asp
 1
<210> 7
<211> 12
<212> PRT
<213> Artificial Sequence
<220>
<223> An artificial peptide
<400> 7
Cys Arg Pro Gly Gln Glu Leu Thr Lys Ser Gly Tyr
                                     10
<210> 8
<211> 24
<212> PRT
<213> Artificial Sequence
```

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<220>
<223> A conserved pattern
<221> SITE
<222> (1) ... (24)
<223> Xaa = Any Amino Acid
<400> 8
Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
                                     10
                                                          15
                  5
 1
Xaa His Xaa Xaa Xaa Cys Xaa Cys
            20
<210> 9
<211> 4
<212> PRT
<213> Mus musculus
<400> 9
Cys Arg Cys Pro
 1
<210> 10
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> A consensus sequence
<221> SITE
<222> (1) ... (4)
<223> Xaa = Any Amino Acid
<400> 10
Cys Xaa Cys Pro
<210> 11
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> A primer
<400> 11
                                                                          25
acctcgaggt cctgtgcatg tgaca
<210> 12
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> A primer
<400> 12
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atgaattctt actgcaggag tgccc

1

25

<210> 13
<211> 11
<212> PRT
<213> Mus musculus

<400> 13
Cys Arg Pro Gly Gln Glu Leu Thr Lys Gln Gly

10